#### **ORIGINAL RESEARCH**



# In vitro cholinesterase inhibitory action of *Cannabis sativa* L. Cannabaceae and in silico study of its selected phytocompounds

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Received: 24 February 2020 / Accepted: 8 January 2021 © The Author(s), under exclusive licence to Springer-Verlag GmbH, DE part of Springer Nature 2021

#### Abstract

*Cannabis sativa* L. Cannabaceae, used for psychoactive rituals in Mesopotamia. Here, we investigated in vitro inhibitory activity of methyl alcohol extract derived from leaves and resin of cannabis against acetylcholinesterase (AChE) and butyryl-cholinesterase (BChE). Moreover, the binding affinity (BA; kcal/mol) of selected phytochemicals of cannabis to AChE and BChE has been predicted in silico. Phytochemicals of cannabis had acceptable BA towards AChE ranging from -6.4 (beta-pinene) to -11.4 (campesterol) and BChE ranging from -5.5 (alpha-pinene) to -9.8 (cannabioxepane). All cannabinoids, flavonoids (apigenin), terpenes, and phytosterols of cannabis were double inhibitors due they utilized hydrogen bonds and hydrophobically interacted with both catalytic triad and peripheral anionic site (PAS) of AChE and BChE. Campesterol is phytosterol docked with AChE and BChE via hydrogen bond and it will be a lead-like molecule for further drug design. Delta-9-Tetrahydrocannabinolic acid has been docked with AChE and BChE and it can be a candidate molecule for further drug design. To sum up, this study not only approved cholinesterase inhibitory effects of cannabis but also suggested an array of phytocompounds as hit small molecules for discovery or design of ecofriendly botanical antiinsectants or phyto-nootropic drugs.

**Keywords** *Cannabis sativa* · Phytonootropics; insecticides · Tetrahydrocannabinolic acid · Terpenes · Phytosterols; cannabinoids

## Introduction

*Cannabis sativa* L. Cannabaceae, hemp, has been used as medicinal herb dates back to 3000 BC (Turner et al. 1980). The seeds and leaves of the plant have therapeutic effects and its psychotropic resins have been used for medical, ritual, or spiritual purposes (Carod-Artal 2013). In Mesopotamia, cradle of civilization, cannabis was utilized for psychoactive or other pharmacological properties (Russo 2007; Teall 2014). It also utilized as insecticide administered orally and topically (McPartland 1997).

Supplementary Information The online version contains supplementary material available at https://doi.org/10.1007/s4020 3-021-00075-0.

☐ Isaac Karimi isaac\_karimi2000@yahoo.com; karimiisaac@razi.ac.ir Cholinesterase (ChE) terminates acetylcholine (ACh) efficacy in synapses (Wang 2013). Two types of ChE reported in vertebrates are known as acetylcholinesterase (AChE; EC 3.1.1.7), and butyrylcholinesterase (BChE; EC 3.1.1.8; Darvesh et al. 2003), found in the brain and in neurofibrillary tangles and neuritic plaques (Beard et al. 1995). Previous studies have shown that the major cause of the cognitive dysfunction in patients with Alzheimer's disease (AD) is a decrease of cholinergic neurotransmission in the cortical regions of the human brain (Schuster et al. 2010). Therefore, these enzymes have been considered as important therapeutic targets for improving the cholinergic deficit which is the main reason for the decline of cognitive and behavioral impairments in AD (Greig et al. 2002).

This study was aimed to investigate the nootropic and/or insecticidal effects of cannabis with a focus on ChE (AChE and BChE) inhibitory effects. In silico molecular docking has been also implemented to elucidate possible mechanisms mediated by main components of cannabis that interact with ChE.

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#### Materials and methods

## **Plant extraction**

The top flowers of *Cannabis sativa* L. have been prepared from local gardens of Iraq in autumn 2018 and authentitized by a botanist in our department while resin (charas = hash-ish = marijuana concentrate) has been bought from herbal store. Specimens were dried for 12 h in a ventilation oven, disassembled, and blended. Then, fine powder (6 g) weighed, homogenized in a glass vial, and extracted with 600 ml of methanol: chloroform (9:1) mixture. After that, the resulting mixture was settled down and centrifuged at 3000 rpm to separate the sediment. The resulting extract has been filtered, diluted, and preserved in a dark glass flask until use.

#### In vitro ChE inhibitory effect

All chemicals were obtained from Sigma-Aldrich (St. Louis, MA, USA) and used as received. For measuring ChE activity (Ellman et al. 1961), the reaction mixture contained 1000 μL Na2HPO<sub>4</sub> buffer (50 mM, pH 7.7), 100 μL of test compound followed by the addition 100 µL of enzyme pre-read at 412 nm using UV-VIS spectrophotometer. Then, contents were pre-incubated for 10 min at 37 °C. The reaction was initiated by the addition 100 µL of substrate (acetylthiocholine iodide and butyrylthiocholine chloride for evaluation of AChE and BChE activities, respectively) and 100 µL of Ellman's reagent (5,5'-dithiobis-(2-nitrobenzoic acid) or DTNB). After 30 min of incubation at 37 °C, absorbance was assayed at 412 nm. All tests were carried out thrice with their respective controls. The percent inhibition was calculated by the help of following equation: Inhibition (%) = (absorbance of negative control—absorbance of test compound)/ (absorbance of negative control)  $\times$  100; where, control = total enzyme activity without inhibitor; test = activity in the presence of test extract. The inhibition percentage of extract has been reported in proportion to eserine as a canonical reversible ChE inhibitor.

#### Molecular docking studies

For docking, crystal structure of protein target was retrieved from Protein Data Bank (PDB) (http://www.RCSB.org; Berman et al. 2002). The PDB format of target proteins, AChE (1EVE: Pacific electric ray (*Torpedo californica*); [Tax Id: 7787] and BChE (1P0I: *Homo sapiens*); [Tax Id:9606] have been edited, optimized and trimmed in Molegro Virtual Docker (Thomsen and Christensen 2006) and Chimera 1.8.1 (http://www.rbvi.ucsf.edu/chimera) before submission to PyRx software version 0.8 (Dallakyan and Olson 2015). The number of each amino acid has been reported in edited 1EVE is Torpedo's number minus one and in edited 1POI is *Homo sapiens*'s number minus three. The structures of the major known bioactive compounds of selected plants were retrieved from PubChem databases (https://pubchem.ncbi. nlm.nih.gov) and ZINC database ver. 12.0 (http://zinc.docki ng.org/) and ChemSpider databases (http://www.chemspider .com). After accomplishment of in silico molecular docking, the results have been presented as binding affinity (BA; kcal/ mol) values. The more negative BA reflects the best pose and binding of ligand to target protein in its binding sites. The selected structure of ligand and target protein were presented here due to limitation of space (vide infra).

#### Pharmacokinetic parameters

The prediction of ADMET (absorption, distribution, metabolism, excretion, and toxicity) properties is an initial phase of drug discovery. The structures of the major phytocompounds were retrieved from aforementioned databases (vide supra) and their canonical SMILES formats submitted to AdmetSAR Database https://lmmd.ecust.edu.Cn/admet sar1 and parameters including blood-brain barrier (BBB) penetration, human gastrointestinal tract (GIT) absorption, distribution, subcellular localization, metabolism via CYP450 2C9, and cardiotoxicity index of *human ether-ago-go-related gene* (HERG) inhibition have been considered as drug-likeness or lead-likeness fitness (Cheng et al. 2012).

# **Results and discussion**

Alzheimer's disease causes loss of intellectual and behavioral abilities and is the most common cause of dementia (Loizzo et al. 2008). Many therapeutic strategies have been developed to treat AD, however ChE inhibitors are forefront. Drugs such as eserine, tacrine, donepezil, rivastigmine, and galanthamine were approved for the treatment of AD, although the search for new inhibitors of AChE and BChE is still continuous (Shoaib et al. 2015).

Enzymologically, AChE and BChE employ an amino acid sequence homology of 65% while they are encoded via various genes located on human chromosomes 7 and 3 (AChE (7q22) and BChE (3q26); Soreq and Zakut 1993). Both AChE and BChE have the same structure (Vellom et al. 1993), where they possess hydrophobic active gorge (Sussman et al. 1991). Upon entering ACh into active site of these enzymes, it binds to two sites including catalytic region adjacent to the base of the gorge and a cholinebinding site midway up gorge. In this line, AChE is specific to hydrolyze ACh whilst BChE has the ability to break many various molecules (Taylor and Radic 1994). The acylation or catalytic site (CS) of AChE composed of Ser200, His440, and Glu327 residues buried at the base of a narrow 20 Å deep gorge, lined predominantly with aromatic residues, whilst catalytic triad in BChE consists of Ser198, His438, and Glu325 (Suárez and Field 2005). The major subsites of AChE are the oxyanion hole (OH) consists of Gly118, Gly119, and Ala201 residues (Ordentlich et al. 1998), while OH site located near of the cholinebinding site and contains Gly116, Gly117 and Ala199 residues in BChE (Bajda et al. 2013). Another site of AChE is anionic subsite (AS) contains aromatic residues (Trp84 and Phe330) and Glu199 in AChE (Kua et al. 2003) whilst AS in BChE contains Trp82, Tyr128, Phe329, and Ala328 residues (Ali et al. 2017). The difference between AChE and BChE lies in the acyl pocket (AP) and the peripheral anionic site (PAS; Houghton et al. 2006). The AP of AChE includes Trp233, Phe288, Phe290, and Phe331 residues (Radic et al. 1992) while AP of BChE includes Leu286 and Val288 residues and PAS of AChE including Tyr70, Asp72, Tyr121, Trp279, and Tyr334 residues (Eichler et al. 1994). The PAS of BChE contains Tyr332 and Asp70 residues (Bajda et al. 2013).

We found that the fraction of resin of *Cannabis sativa* L. (hashish) showed inhibitory effects of 80.00% and 68.00% for AChE and BCHE, respectively (p=0.001). While fraction of leaves of Cannabis sativa L. (marihuana) showed similar inhibitory effects of 52.33% and 49.00% for AChE and BCHE, respectively (p=0.226). Moreover, phytocompounds of Cannabis sativa L. that showed acceptable BA with AChE and BChE have been reported at this juncture. In this continuum, delta-3-carene or 3-carene (Fig. 1) is a bicyclic monoterpenoid alkene found naturally in cannabis and white pepper (Kendall and Alexander 2017), has AChE inhibitory effects (Lomarat et al. 2015). The 3-carene has been docked with AChE with BA of -7.3 kcal/mol. The 3-carene has hydrophobic interactions with amino acid residues in the catalytic triad (Trp83, Phe329 and Tyr333) and PAS (Tyr120) and AS (Phe330; Supplementary material (SM)). In this line, it has appropriate BA (-6.1 kcal/ mol) with BChE via hydrophobic interactions with amino acid residues of PAS (Tyr329) and AS (Ala325, Trp79 and Phe326) and the CS (His433; SM). ADMET properties of 3-carene showed that it can go through BBB and absorbed from GIT whilst it is a weak inhibitor of HERG (Table 3).

Agarospirol is a sesquiterpene found in cannabis (Rothschild et al. 2005). This study showed that agarospirol has been docked with AChE (BA of –8 kcal/mol) via hydrogen bonds with hydroxyl group of Ser121 and through hydrophobic interactions with PAS (Tyr333, Tyr120 and Asp71) and AS (Trp83 and Phe329) and AP (Phe330) and with His439 residue of CS (SM). It also docked with BChE through hydrophobic interactions with amino acid residues in AS (Trp79) and PAS (Asp67 and Tyr329) and His433 residue of the CS (SM). According to study ADMET properties, it can be absorbed from BBB and GIT, while localized in lysosome (Table 3).

Alpha-bisabolene is a sesquiterpene found in cannabis (Gulluni et al. 2018). In this study, it has been docked with AChE with BA of –9.2 kcal/mol via hydrophobic interactions with amino acid residues in AP (Phe330) and PAS (Tyr333 and Tyr120) and AS (Trp83 and Phe329) and OH (Gly117) and CS (His439; SM). In this regard, alphabisabolene has been docked with BChE with appropriate BA (Table 2) through hydrophobic interactions with amino acid residues in PAS (Tyr329) and AS (Ala325, Phe326, Trp79 and Tyr125; SM). Based on ADMET results, it can be absorbed from GIT and crossed BBB (Table 3).

Alpha-bisabolol is a sesquiterpene found in cannabis (Rothschild et al. 2005) and several plants. It is a strong inhibitor of AChE (Shanmuganathan et al. 2015). In this study, alpha-bisabolol has been docked with AChE with BA (-8.7 kcal/mol; Table1) through hydrogen bond with Asp71 residue of PAS and through hydrophobic interactions with Trp83, Phe329 and Glu198 residues of AS and Tyr120 and Tyr333 residues of PAS and Phe330 residues of AP and His residues of CS. Alpha-bisabolol has been docked with BChE (BA -7.5 kcal/mol; Table 2) through hydrophobic interactions with amino acid residues in PAS (Tyr329 and Asp67) and AS (Ala325, Phe326, Trp79 and Tyr125) and the CS (His433; SM). The ADMET assays showed that it can be absorbed orally and transferred through BBB while it is a weak inhibitor of HERG (Table 3).

Alpha-guaiene is a sesquiterpene found in many plants including cannabis (Fischedick et al. 2010) and works as an insect repellent (Murugesan et al. 2012). This study showed, alpha-guaiene has been docked with AChE (-8.9 kcal/mol; Table1) via hydrophobic interactions with amino acid residues of PAS (Tyr333, Tyr120 and Asp71) and AP (Phe330) and catalytic AS (Trp83 and phe330; SM). In addition, it hydrophobically interacts with amino acid residues in PAS (Tyr329 and Asp67) and AS (Phe326 and Trp79) and CS (His433) of BChE (SM) with BA -7.3 kcal/mol (Table 2). The ADMET properties results of alpha-guaiene showed that it can be transferred via BBB and absorbed from GIT; it is a weak inhibitor of HERG (Table 3).

Alpha-humulene is a sesquiterpene found in several medicinal plants like cannabis (Fischedick et al. 2010), *Commiphora myrrha* (Nees), Engler and ginseng (Gitau 2015). Alpha-humulene contained in herbal extracts have insecticidal effects through inhibition of AChE (Lee and Ahn 2013). In this study, alpha-humulene has been docked with AChE with acceptable BA (-9.0 kcal/mol) through hydrophobic interactions with amino acid residues in PAS (Tyr120 and Asp71) and catalytic triad (Trp83; SM). Alpha-humulene has been hydrophobically docked with BA of -7.7 kcal/mol. In this regard, alpha-humulene interacted with amino acid residue in catalytic triad (His433) and PAS

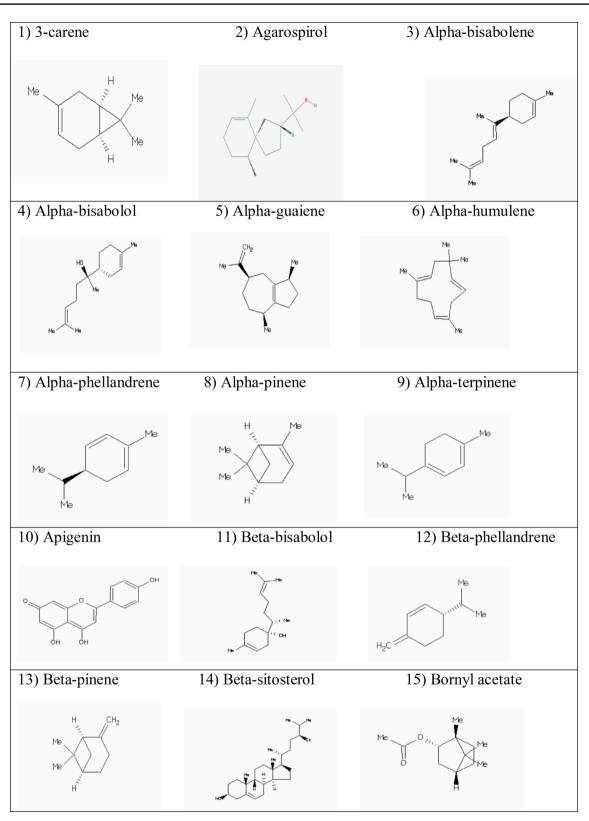


Fig. 1 The selected phytochemicals of Cannabis sativa L

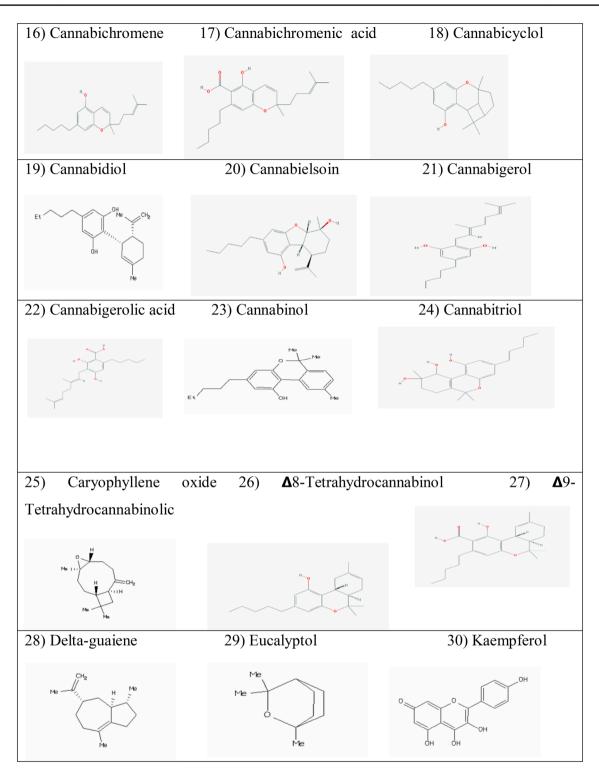
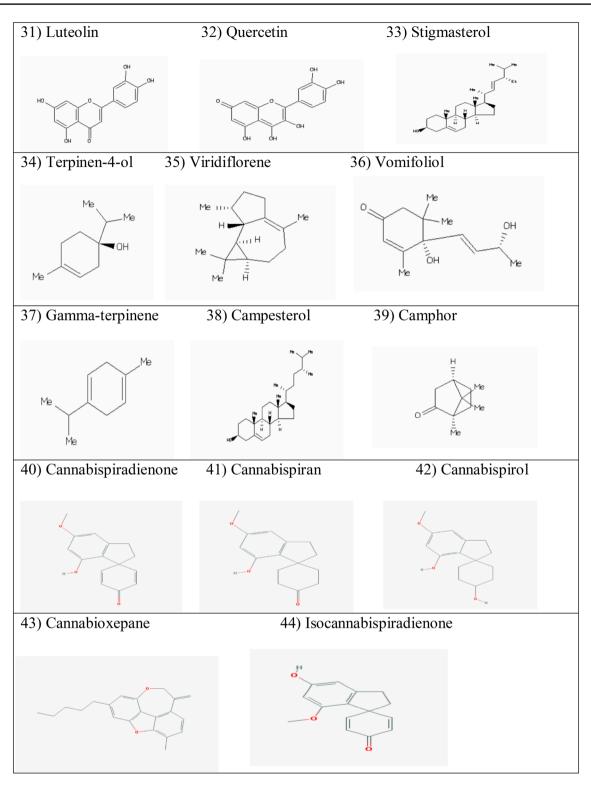


Fig. 1 (continued)

(Tyr329 and Asp67) and AS (Phe326, Trp79 and Ala325) of BChE. Based on ADMET properties, it can pass BBB and GIT (Table 3).

Alpha and beta-phellandrene are monoterpenes found in cannabis (Fischedick et al. 2010). Alpha-phellandrene has insecticidal and pesticidal activity (Onyenekwe et al. 2012). In this study, alpha-phellandrene interacted hydro-phobically with amino acid residues in PAS (Tyr120, Tyr333 and Asp71), AS (Phe329), AP (Phe330) and CS (His439; SM) of AChE (BA of -6.6 kcal/mol). Beta-phellandrene



#### Fig. 1 (continued)

also interacted hydrophobically with amino acid residues in PAS (Tyr333), AS (Trp83 and Phe329), AP (Phe330) and CS (His439; SMSM) of AChE (BA of -7 kcal/mol). Alpha-phellandrene has been loosely docked with BChE (BA - 5.8 kcal/mol; Table 2). While beta-phellandrene has been docked with BChE (BA - 6 kcal/mol; Table 2) via hydrophobic interactions with amino acid residues of AS (Tyr125, Trp79 and Ala325) and CS (His433; SM). The

Table 1 In silico molecular
docking of major
phytochemicals of Cannabis
sativa L. against AChE (PDB
code 1EVE)

Ligand	Binding affinity	RMSD/UB	RMSD/LB
3-Carene	-7.3	5.115	2.521
Agarospirol	-8	3.392	1.753
Alpha-bisabolene	-9.2	6.7	1.296
Alpha-bisabolol	-8.7	2.305	1.905
Alpha-guaiene	-8.9	5.216	1.658
Alpha-humulene	-8.9	4.833	1.533
Alpha-phellandrene	-6.6	5.105	2.371
Alpha-pinene	-6.6	4.202	1.947
Alpha-terpinene	-7.4	4.25	0.964
Apigenin	-9.9	5.793	2.522
Beta-bisabolol	-8.3	2.712	1.55
Beta-phellandrene	-7	4.116	1.507
Beta-pinene	-6.4	3.376	1.016
Beta-Sitosterol	-10.9	2.504	1.353
Bornyl Acetate	-7.1	2.89	1.928
Campesterol	-11.4	3.222	1.703
Camphor	-6.6	2.695	1.505
Cannabichromene	-9.4	7.731	1.913
Cannabichromenic	-9.3	9.687	3.726
Cannabicyclol	-9.6	8.896	3.467
Cannabidiol	-9.8	1.778	0.315
Cannabielsoin	-8.9	4.138	2.003
Cannabigerol	-8.9	2.51	0.907
Cannabigerolic acid	-9.2	9.104	2.069
Cannabinol	-10.1	3.461	1.348
Cannabitriol	-9.8	7.238	2.23
Caryophyllene oxide	-9.0	4.5	2.045
Delta8-Tetrahydrocannabinol	-10.1	6.03	1.86
Delta9-Tetrahydrocannabinolic acid	-10.3	8.8	3.978
Delta-guaiene-85598968	-8.8	4.915	1.32
Eucalyptol	-6.7	2.22	1.432
Gamma-Terpinene	-7.4	4.314	0.922
Kaempferol	-10.1	5.646	2.428
Luteolin-18185774-Mh	-9.9	2.662	1.143
Quercetin	-9.8	4.156	2.069
Stigmasterol	-11.3	1.196	0.986
Terpinene-4-Ol	-7.1	4.513	1.104
Viridiflorene	-8.7	6.398	3.745
Vomifoliol	-7.8	5.191	3.09
Cannabispiradienone	-9.2	6.772	2.527
Cannabispiran	-8.9	3.009	2.019
Cannabispirol	-9.2	1.595	1.375
Cannabioxepane	-10.4	7.894	2.685
Isocannabispiradienone	-9.4	7.727	5.157

RMSD root mean-square deviation is the measure of the average distance between the atoms, UB upper bound, LB lower bound

ADMET assays showed that alpha and beta-phellandrene can be absorbed orally and crossed BBB while they are week inhibitor of HERG (Table 3).

Alpha and beta-pinene are monoterpenes. Pinene has capability to cross the BBB readily where it can inhibit activity of ACHE, leading to better memory results (Owokotomo

Table 2 In silico molecular docking of major phytochemicals of Cannabis sativa L. against BChE (PDB code 1P0I)

Ligand	Binding affinity	RMSD/UB	RMSD/LB 1.536
3-carene	-6.1	2.671	
Agarospirol	-7.4	2.667	1.513
Alpha-bisabolene	-7.1	6.57	1.11
Alpha-bisabolol	-7.5	1.891	1.193
Alpha-guaiene	-7.3	5.44	2.671
Alpha-humulene	-7.7	4.411	1.365
Alpha-phellandrene	-5.8	21.44	20.187
Alpha-pinene	-5.5	3.404	0.963
Alpha-terpinene	-6.1	2.279	1.131
Apigenin	-9.1	5.312	2.857
Beta-bisabolol	-6.9	6.161	1.235
Beta-phellandrene	-6.0	2.42	1.438
Beta-pinene	-5.6	20.444	19.181
Beta-sitosterol	-9.3	8.016	3.152
Bornyl acetate	-6.3	4.046	2.144
Campesterol	-9.3	8.235	3.201
Camphor	-6.6	2.695	1.505
Cannabichromene	-7.7	21.782	19.152
Cannabichromenic	-7.7	4.05	3.681
Cannabicyclol	-8.4	4.604	3.004
Cannabidiol	-8.2	3.316	1.856
Cannabielsoin	-8.9	3.631	1.341
Cannabigerol	-7.3	5.47	3.775
Cannabigerolic acid	-7.5	19.972	17.886
Cannabinol	-8.8	4.942	3.402
Cannabitriol	-8.7	7.986	4.66
Caryophyllene oxide	-7.4	4.192	2.067
Delta8-Tetrahydrocannabinol	-8.7	5.913	2.114
Delta9-Tetrahydrocannabinol		5.861	2.358
Delta-Guaiene-85598968	-7.5	4.023	1.143
Eucalyptol	-5.7	21.312	19.967
Gamma-terpinene	-5.7	2.965	1.755
Kaempferol	-8.5	5.679	4.208
Luteolin-18185774-mh	-9.4	2.76	1.074
Quercetin	-9	6.803	5.141
Stigmasterol	-9.7	2.594	1.798
Terpinene-4-ol	-5.9	4.512	2.743
Viridiflorene	-8.0	3.884	1.355
Vomifoliol	-6.7	20.743	19.476
Cannabispiradienone	-8.2	5.875	2.186
Cannabispiran	-8.2	5.942	2.23
Cannabispirol	-8.1	6.517	3.638
Cannabioxepane	-9.8	3.235	0.658
Isocannabispiradienone	-8.3	4.914	2.007

*RMSD* root mean-square deviation is the measure of the average distance between the atoms, *UB* upper bound, *LB* lower bound

et al. 2015). Alpha-pinene has insecticidal and pesticidal activities (Onyenekwe et al. 2012). This study showed that alpha and beta-pinene have been docked with AChE with

BA of -6.6 and -6.4 kcal/mol, respectively. In this line, alpha-pinene interacted hydrophobically with amino acid residues in PAS (Tyr333, Tyr120) and AP (Phe330) and CS

### Table 3 In silico ADMET properties of phytochemicals of Cannabis sativa L. by admetSAR

Compound	BBB	HIA	CYP inhibition/substrate	Subcellular localization	HERG Inhibition
3-Carene	0.9473	0.9943	Non-substrate/Non-inhibitor	Lysosome	Weak
Agarospirol	0.9669	0.9954	Non-substrate/Non-inhibitor	Lysosome	Weak
Alpha-bisabolene	0.9399	0.9934	Non-substrate/Non-inhibitor	Lysosome	Weak
Alpha-bisabolol	0.9667	0.9900	Non-substrate/Non-inhibitor	Mitochondria	Weak
Alpha-guaiene	0.9723	0.9940	Non-substrate/Non-inhibitor	Lysosome	Weak
Alpha-humulene	0.9733	0.9972	Non-substrate/Non-inhibitor	Lysosome	Weak
Alpha-phellandrene	0.9049	0.9965	Non-substrate/Non-inhibitor	Lysosome	Weak
Alpha-pinene	0.8959	0.9964	Non-substrate/Non-inhibitor	Lysosome	Weak
Alpha-terpinene	0.9415	0.9953	Non-substrate/Non-inhibitor	Nucleus	Weak
Apigenin	0.6364	0.9887	Non-substrate/inhibitor	Mitochondria	Weak
Beta-bisabolol	0.9462	0.9899	Non-substrate/Non-inhibitor	Mitochondria	Weak
Beta-phellandrene	0.9261	0.9830	Non-substrate/Non-inhibitor	Lysosome	Weak
Beta-pinene	0.9229	0.9834	Non-substrate/Non-inhibitor	Lysosome	Weak
Beta-sitosterol	0.9743	1.0000	Non-substrate/Non-inhibitor	Lysosome	Weak
Bornyl acetate	0.9719	0.9969	Non-substrate/ Non-inhibitor	Mitochondria	Weak
Campesterol	0.9749	1.0000	Non-substrate/Non-inhibitor	Mitochondria	Weak
Camphor	0.9836	0.9971	Non-substrate/inhibitor	Mitochondria	Weak
Cannabichromene	0.9672	0.9937	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabichromenic acid	0.5785	0.8983	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabicyclol	0.9791	0.9926	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabidiol	0.6597	0.9937	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabielsoin	0.9379	0.9964	Non-substrate/inhibitor	Mitochondria	Weak
Cannabigerol	0.5888	0.9960	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabigerolic acid	0.6891	0.9773	Non-substrate/inhibitor	Mitochondria	Weak
Cannabinol	0.9856	0.9922	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabitriol	0.7859	0.9707	Non-substrate/Non-inhibitor	Mitochondria	Weak
Caryophyllene oxide	0.9474	0.9950	Non-substrate/inhibitor	Lysosome	Weak
$\Delta^8$ -THC	0.9685	0.9949	Non-substrate/ inhibitor	Mitochondria	Weak
$\Delta^9$ -THC	0.5953	0.9164	Non-substrate/inhibitor	Mitochondria	Weak
Delta-guaiene	0.9723	0.9940	Non-substrate/Non-inhibitor	Lysosome	Weak
Eucalyptol	0.9842	0.9880	Non-substrate/Non-inhibitor	Lysosome	Weak
Gamma-terpinene	0.9431	0.9972	Non-substrate/Non-inhibitor	Lysosome	Weak
Kaempferol	0.6286	0.9855	Non-substrate/inhibitor	Mitochondria	Weak
Luteolin	0.5711	0.9650	Non-substrate/Non-inhibitor	Mitochondria	Weak
Quercetin	0.5711	0.9650	Non-substrate/Non-inhibitor	Mitochondria	Weak
Stigmasterol	0.9743	1.0000	Non-substrate/Non-inhibitor	Lysosome	Weak
Terpinene-4-Ol	0.9104	0.9969	Non-substrate/Non-inhibitor	Mitochondria	Weak
Viridiflorene	0.9790	1.0000	Non-substrate/Non-inhibitor	Lysosome	Weak
Vomifoliol	0.5351	1.0000	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabispiradienone	0.9259	1.0000	Non-substrate/inhibitor	Mitochondria	Weak
Cannabispiran	0.9225	0.9967	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabispirol	0.8235	0.9957	Non-substrate/Non-inhibitor	Mitochondria	Weak
Cannabioxepane	0.9709	1.0000	Non-substrate/Non-inhibitor	Plasma membrane	Weak
Isocannabispiradie-	0.9259	1.0000	Non-substrate/Non-inhibitor	Mitochondria	Weak

ADMET absorption, distribution, metabolism, and excretion-toxicity, BBB blood-brain barrier penetration, HIA human intestinal absorption, CYP cytochrome P, HERG human ether-a-go-go-related genes inhibition

(His439) of AChE while beta-pinene interacted hydrophobically with amino acid residues of AS (Trp83 and Phe329) and AP (Phe330) solely with His439 residues in CS. Alpha and beta-pinene have been hydrophobically docked with BChE with BA of -5.5 and -5.6 kcal/mol, respectively. In this line, beta-pinene has been docked with BChE through hydrophobic interactions with amino acid residues of PAS (Tyr329) and AS (Phe326, Trp79 and Ala325; SM). Based on ADMET properties, they can be absorbed orally and passed through BBB without considerable side effects (Table 3).

Alpha and gamma-terpinene are monoterpenes found in an array of medicinal plants like cannabis and tea tree oil (Miyazawa and Yamafuji 2006; Hartsel et al. 2016). In the pioneered work, it has been reported that herbal extracts containing terpinene possess inhibitory effects of AChE (Miyazawa and Yamafuji 2006). Alpha-terpinene has hydrophobic interactions with amino acid residues in PAS (Tyr333) and AS (Trp83 and Phe329) and AP (Phe330). Gamma-terpinene has also hydrophobic interactions with amino acid residues in PAS (Tyr333), AS (Trp83 and Phe329), CS (His439) and AP (Phe330; SM). Alpha and gamma-terpinene have been hydrophobically docked with BChE with BA of -6.1 and -5.7 kcal/mol, respectively. In this line, alpha-terpinene has been docked with BChE through hydrophobic interactions with amino acid residues of PAS (Tyr329) and AS (Phe326, Tyr125, Trp79 and Ala325) and CS (His433). Gamma-terpinene has hydrophobic interactions with amino acid residues in AS (Tyr125, Trp79 and Ala325) and CS (His433; SM). The ADMET prediction showed that they can be crossed from GIT and BBB (Table 3).

Apigenin is a flavonoid compound found in cannabis (Hartsel et al. 2016). Apigenin has ability to inhibit AChE activity (Balkis et al. 2015). Apigenin has been docked with AChE with BA –9.9 kcal/mol via hydrogen bonds with amino acid residues in AS (Glu198) and with Tyr69 in PAS and with hydroxyl group of Gln68. It also has hydrophobic interactions with amino acid residues of PAS (Asp71 and Tyr120), AS (Trp83 and Phe329) and CS (His439) of AChE. The Tyr125 residue of PAS and Ala325 residue of AS interacted with apigenin with hydrogen bond while Trp79 residue of AS interacted hydrophobically with apigenin (SM). Based on ADMET, it can be absorbed from GIT, crossed BBB, and localized in mitochondria while it is a week inhibitor of HERG (Table 3).

Beta-bisabolol is a sesquiterpene found in cannabis (Hartsel et al. 2016). In present study, beta-bisabolol has been docked with AChE with BA (-8.3 kcal/mol; Table) via hydrogen bonding with Tyr120 and several hydrophobic interactions with amino acid residues of PAS (Tyr333 and Asp71), AS (Trp83 and Phe329), AP (Phe289 and Phe330) and CS (His439). Beta-bisabolol docked with BChE with BA -6.9 kcal/mol (Table 2) through hydrophobic interactions with amino acid residues in PAS (Tyr329 and Asp67) and AS (Ala325, Phe326, Trp79 and Tyr125) and CS (His433; SM). Based on ADMET properties, it has ability to penetrate BBB and cross GIT while it weak HERG inhibitor (Table 3).

Beta-sitosterol is a phytosterol found in cannabis (Hartsel et al. 2016). Beta-sitosterol has been docked with AChE (BA -10.9 kcal/mol) via hydrophobic interactions with amino acid residues in PAS (Tyr333, Asp71, Tyr120 and Trp278), AS (Trp83 and Phe329) and AP (Phe289, Phe330 and Phe289; SM). It has been docked with BChE with BA of -9.3 kcal/mol through hydrophobic interactions with amino acid residues in AS (Trp79, Phe326 and Ala325) and PAS (Tyr329 and Asp67) and CS (His 433; SM). Beta-sitosterol can be absorbed from GIT, crossed BBB, and localized in lysosome while it is a week inhibitor of HERG (Table 3).

Bornyl acetate is a monoterpene found in cannabis (Hartsel et al. 2016). Bornyl acetate has insecticidal activities against several warehouse insects and mites (Li et al. 2015). In this study, bornyl acetate has a BA of -7.1 kcal/mol with AChE through two hydrogen bonds with amino acid Gly117 and Gly118 in OH and with amino acid Ser199 in CS (SM) and via hydrophobic interactions with Tyr120 residues of PAS and with amino acid residues of AS (Phe329 and Trp83) and AP (Phe330; SM). Bornyl acetate has been docked with BChE (BA of -6.3 kcal/mol) through hydrophobic interactions with amino acid residues in AS (Trp79, Phe326 and Ala325) and PAS (Tyr329) and CS (His 433; SM). Based on ADMET results, it can be absorbed from GIT and crossed BBB (Table 3).

Campesterol is another phytosterol found in cannabis (Montserrat-de la Paz et al. 2014). Campesterol has cholinesterase inhibitory activity (Elufioye et al. 2017). In this study, campesterol has been docked with AChE (BA of -11.4 kcal/mol) via hydrogen bonds with hydroxyl group of Ser285 and through hydrophobic interactions with OH (Gly 117), AP (Phe330 and Phe 289), AS (Trp83, Phe329 and Glu198) and PAS (Tyr333, Tyr120 and Asp71). Campesterol has been docked with BChE (BA of -9.3 kcal/mol) through hydrogen bonds with hydroxyl group of Ser284 and through hydrophobic interactions with PAS (Asp67 and Tyr329) and AS (Ala325 and Trp79) and CS (His433; SM). Campesterol can be absorbed from GIT, crossed BBB, and localized in mitochondria while it is a week inhibitor of HERG (Table 3).

Camphor is a monoterpene found in cannabis (Fischedick et al. 2010). In this study, Ser199 as crucial amino acid residue in CS of AChE interacted with camphor through hydrogen binding with length 3.33 Å while Gly117, His439, Phe 330 and Trp83 residues in catalytic triad interacted hydrophobically with camphor. Camphor has a BA of -5.9 kcal/mol with BChE through hydrogen bond with amino acid

Tyr329 in PAS with a bond length of 2.75 Å (SM) and through hydrophobic interactions with (Ala325, Phe326and Trp79) residues of AS. Camphor can be absorbed from GIT, crossed BBB, and localized in mitochondria while it is a week inhibitor of HERG (Table 3).

Phytocannabinoids found in cannabis including cannabidiol, cannabielsoin, cannabigerol, cannabigerolic acid, cannabinol and cannabitriol (Hartsel et al. 2016) have been docked with AChE and BChE with satisfactory BA. In this line, cannabidiol docked with AChE via hydrophobic interactions with PAS (Tyr333, Trp278, Asp71 and Tyr120) and AS (Trp83 and Phe329) AP (Phe289 and Phe330). Cannabidiol has been docked with BChE with BA of –8.2 kcal/ mol through hydrogen bonds with hydroxyl group of Thr117 and through hydrophobic interactions with AS (Trp79 and Phe 326) and AP (Leu283) and CS (His433 and Ser195; SM).

The Asp71 residue of PAS interacted with cannabielsoin with hydrogen bond while Tyr333, Tyr120, Trp69 and Trp278 residues of PAS and Phe330, Phe287 residues of AP and Phe329 and Trp83 residues of AS interacted hydrophobically with cannabielsoin (SM). Cannabielsoin docked with BChE via hydrogen bonding with Trp79 and hydroxyl group of Gly75 and with several hydrophobic interactions with amino acid residues of PAS (Tyr329 and Asp67) and AS (Ala325) and CS (His433; SM).

Cannabigerol interacted with AChE via hydrogen bond with Tyr120 and with several hydrophobic interactions with amino acid residues of PAS (Asp71, Tyr333 and Trp278) and AS (Phe329, Glu198 and Trp83) and AP (Phe330; SM). Cannabigerol has a BA of -7.3 kcal/mol with BChE through hydrogen bonds with Trp79 of AS and hydroxyl group of Asn80 (SM) and through hydrophobic interactions with Asp67 of PAS and Tyr125 of AS and His433 of CS.

Cannabigerolic docked with AChE through two hydrogen bonds with amino acid Tyr120 and one with Asp71 in PAS (SM) and through hydrophobic interactions with (Tyr333, Trp69 and Trp278) and AS (Trp83 and Phe329) and AP (Phe287 and Phe330; Figure). In this regard, cannabigerol has been docked with BChE (BA – 7.5 kcal/mol) through hydrogen bonds with hydroxyl group of Ser284 and through hydrophobic interactions with AS (Ala325, Trp79 and Phe326) and PAS (Tyr329 and Asp67) and AP (Leu283) and CS (His433; SM).

Cannabinol also docked with AChE through hydrogen bond with amino acid Tyr120 in PAS with a bond length of 2.75 (SM) and through hydrophobic interactions with several amino acid residues of activity site of AChE. Cannabinol docked with BChE (BA of -8.8 kcal/mol) through hydrogen bonds with hydroxyl group of Thr117 and through hydrophobic interactions with PAS (Asp67 and Tyr329) and AS (Ala325, Tyr125, Trp79 and Phe326) and CS (His433; SM). Cannabitriol docked with AChE through hydrogen bond with amino acid Tyr120 and with through hydrophobic interactions with several amino acid residues in PAS (Tyr333, Trp69 and Trp278) and AS (Phe329 and Trp83) and AP (Phe330; SM). Cannabitriol has been docked with BChE (BA of -8.7 kcal/mol) through hydrogen bonds with His433 and through hydrophobic interactions with CS (Ser195) and PAS (Tyr329) and AS (Ala325, Trp79 and Phe326). All phytocannabinoids have accepted ADMET properties to be considered as lead-like compounds.

Cannabichromene is a cannabinoid found in cannabis (Hartsel et al. 2016). In present study, cannabichromene has been docked with AChE with BA of –9.4 kcal/mol via hydrogen bonds with Asp71 and Tyr120 residues and hydrophobic interactions with amino acid residue of PAS (Tyr333) and AS (Trp83 and Phe329) and AP (Phe330) and OH (Gly117; SM). Cannabichromene has been docked with BChE via hydrogen bonds with hydroxyl group of Thr117 and through hydrophobic interactions with AS (Ala325, Trp79 and Phe326) and PAS (Asp67 and Tyr329) and AP (Leu283) and CS (His433 and Ser195; SM). The ADMET properties assay showed that cannabichromene can be absorbed orally and passed through BBB without considerable side effects.

Cannabichromenic acid is a cannabinoid found in cannabis (Hartsel et al. 2016). In present study, cannabichromenic acid has BA –9.3 kcal/mol with AChE through hydrogen bonds with amino acid residue of AP (Phe287) and with hydroxyl group of Arg288 and through hydrophobic interactions with PAS (Tyr333, Asp71, Trp278 and Tyr120) and AP (Phe330) and AS (Trp83 and Phe329; SM). Cannabichromenic acid has been docked with BChE with BA -7.7 kcal/mol via hydrophobic interactions with amino acid residues of PAS (Asp67 and Tyr329) and AS (Ala325, Trp79 and Phe326) and CS (His433; SM). The ADMET properties showed cannabichromenic acid can be crossed BBB and GIT and distributed into mitochondria (Table 3).

Cannabicyclol is other cannabinoid found in cannabis (Hartsel et al. 2016). In this study, it has been docked with AChE with BA of –9.6 kcal/mol through hydrophobic interactions with amino acid residues in AP (Phe289 and Phe330) and PAS (Tyr333, Asp71 and Tyr120) and CS (His439) and AS (Phe329 and Trp83; SM). Cannabicyclol has a BA of –8.4 kcal/mol with BChE through two hydrogen bonds with hydroxyl group of Thr117 and through hydrophobic interactions with PAS (Asp67 and Tyr329) and AP (Leu283 and Val285) and AS (Ala325, Tyr125, Trp79 and Phe326) and CS (His433; SM). The ADMET properties of cannabicyclol showed that it can be transferred through BBB and orally absorbed from GIT while it is a weak inhibitor of HERG (Table 3).

Caryophyllene oxide is a sesquiterpene found in cannabis (Hartsel et al. 2016). Caryophyllene oxide has been docked

with BA of -9 kcal/mol via several hydrophobic interactions with PAS (Tyr120, Tyr278, Tyr 333, Tyr69 and Asp71) and AS (Trp83 and Phe329) and AP (Phe330) and CS (His439; SM). Asp67 and Tyr329 residues of PAS and Ala325, Trp79 and Phe326 residues of AS interacted hydrophobically with caryophyllene oxide (SM). Based on ADMET properties, caryophyllene oxide can be absorbed from GIT and crossed BBB (Table 3).

 $\Delta^8$ -Tetrahydrocannabinol ( $\Delta^8$ -THC) and  $\Delta^9$ -Tetrahydrocannabinol ( $\Delta^9$ –THC) are phytocannabinoids (Hartsel et al. 2016). In this study,  $\Delta^8$ -THC has been hydrophobically docked with amino acid residues in PAS (Tyr120, Asp71 and Tyr333) and AS (Trp83 and Phe329) and AP (Phe330, Phe289) and OH (Gly117) of AChE (SM) with appropriate BA of -10.1 kcal/mol, while  $\Delta^9$ -THC has been docked with AChE with BA -10.3 kcal/mol via hydrogen bonding with Tyr120 and several hydrophobic interactions with amino acid residues of PAS (Tyr333, Tyr69, Trp278 and Asp71) and AP (Phe330 and Phe287) and AS (Trp83 and Phe329; SM). In this line,  $\Delta^8$ -THC has been docked with BChE (-8.7 kcal/mol) through hydrophobic interactions with AS (Ala325, Tyr125, Trp79 and Phe326) and PAS (Tyr329), while  $\Delta$ 9-THC hydrophobically interaction with PAS (Asp67) and AP (Leu283) and AS (Ala325, Tyr125, Trp79 and Phe326) and CS (His433 and Ser195) of BChE (SM). Based on ADMET results, THC can be absorbed from GIT and crossed BBB, localized in mitochondria, while it is a weak inhibition of HERG (Table 3).

Delta-guaiene is a sesquiterpene found in cannabis (Romano and Hazekamp 2013). In present study, deltaguaiene has been docked with AChE (BA -8.8 kcal/mol) via hydrophobic interactions by Asp71, Tyr120 and Tyr333 in PAS and by Trp83 and Phe329 in AS and by Phe330 in AP (SM). In addition, it docked with BChE with BA -7.5 kcal/ mol through hydrophobic interactions with AS (Ala325, Trp79 and Phe326) and PAS (Tyr329) and CS (His433; SM). Delta-guaiene can be absorbed from GIT, crossed BBB, and localized in lysosome while it is a week inhibitor of HERG (Table 3).

Eucalyptol is a monoterpene found in cannabis (Hartsel et al. 2016). It used as insecticide and insect repellent (Klocke et al. 1987; Sfara et al. 2009). In this study, eucalyptol has been docked with AChE via hydrogen bonds with Tyr120 and through hydrophobic interactions with Tyr333 in PAS and Phe329 in AS and Phe330 in AP and Gly117 in OH (SM). In this regard, eucalyptol has been docked with BChE (-5.7 kcal/mol) through hydrophobic interactions (SM). The ADMET properties of eucalyptol showed that it can be transferred via BBB and absorbed from GIT while it tends to be localized in lysosome (Table 3).

Kaempferol is a flavonoid found in cannabis (Flores-Sanchez and Verpoorte, 2008). In a seminal work, kaempferol has showed AChE inhibitory potential (Ding et al. 2013). In this study, kaempferol has been docked with AChE (BA -10.1 kcal/mol; Table). The Tyr69 residue of PAS and Glu198 residue of AS interacted with kaempferol with hydrogen bonds while Tyr120 and Asp71 residues of PAS and Phe329 and Trp83 residues of AS interacted hydrophobically with kaempferol (SM). Kaempferol has been docked with BChE with BA -8.5 kcal/mol via hydrogen bonding with Tyr329 and Asp67 residues of PAS and with His433 residue of CS and with hydroxyl group of Asn65 (SM) and through hydrophobic interactions with amino acid residues of AS (Trp79). The ADMET properties showed kaempferol can be crossed BBB and GIT and distributed into mitochondria (Table 3).

Luteolin is a flavonoid found in cannabis (Flores-Sanchez and Verpoorte, 2008). Luteolin induces neurite outgrowth and enhanced cholinergic activities (Omri et al. 2012). In this study, luteolin has a BA of -9.9 kcal/mol with AChE through two hydrogen bonds with Glu198 in AS and Tyr69 in PAS and with hydroxyl group of Gln68 (SM) and through hydrophobic interactions with amino acid residues in PAS (Tyr120 and Asp71) and AS (Trp83 and Phe329). In this line, luteolin has a BA -9.4 kcal/mol with BChE via hydrogen bonding with PAS (Asp67) and CS (His433) and two hydroxyl group of Asn65 and Asn80 (SM) and through hydrophobic interactions with AS (Trp79). Based on ADMET properties, luteolin has ability to penetrate BBB and cross GIT while it weak HERG inhibitor (Table 3).

Quercetin is another flavonoid found in cannabis (Flores-Sanchez and Verpoorte 2008). Quercetin exerts the cognitive enhancing effect in rat model of Parkinson's disease via its antioxidative effect resulting in the promotion of neuron survival (Sriraksa et al. 2012). Quercetin has been docked with AChE (-9.8 kcal/mol) through hydrogen bonds with Glu198 in AS and Tyr69 in PAS and with hydroxyl group of Gli68 (SM) and through several hydrophobic interactions with amino acid residues in PAS (Tyr120 and Asp71) and AS (Trp83 and Phe329). Quercetin has been docked with BChE with BA -9 kcal/mol via hydrogen bonding with PAS (Tyr329 and Asp67) and CS (His433) and with two hydroxyl group of Asn65 and Glu194 (SM) and through hydrophobic interactions with Trp79 residue of AS. Based on ADMET properties, quercetin has ability to pass BBB and GIT (Table 3).

Stigmasterol is a phytosterol found in cannabis (Ryz et al. 2017). In this study, stigmasterol has been docked with AChE with BA of -9.7 kcal/mol through hydrophobic interactions with AP (Phe289) and PAS (Tyr333, Trp278 and Tyr120) and AS (Phe329 and Trp83; SM). Stigmasterol has been docked with BChE (BA of -11.3 kcal/mol) through hydrogen bonds with two hydroxyl groups of Asn286 and Ser284 and through hydrophobic interactions with PAS (Tyr329 and Asp67) and AS (Trp79 and Phe326) and CS (His433; SM). The ADMET properties of stigmasterol

showed that it can be absorbed from GIT and transferred through BBB while it tends to be localized in lysosome (Table 3).

Terpinen-4-ol is a terpene found in cannabis (Grotenhermen and Russo 2006). It has the ability to inhibit AChE (Dohi et al. 2009). In this study, terpinen-4-ol has a BA of -7.1 kcal/mol with AChE through hydrogen bonds with Tyr120 residue in PAS and through hydrophobic interactions with Tyr333 residue of PAS and Phe330 of AP and Trp83 of AS (SM). In addition, it has been docked with BChE with BA of -5.9 kcal/mol through hydrophobic interactions with PAS (Tyr329) and AS (Trp79) and CS (His433; SM). Based on ADMET properties, terpinen-4-ol has ability to penetrate BBB and cross GIT while it weak HERG inhibitor (Table 3).

Viridiflorene is a sesquiterpene found in cannabis (Hartsel et al. 2016). This study showed that viridiflorene has been docked with AChE with BA of -8.7 kcal/mol through hydrophobic interactions with PAS (Tyr333, Tyr120 and Asp71) and AS (Trp83 and Phe329) and AP (Phe330; SM). Viridiflorene has a BA of -8 kcal/mol with BChE through hydrophobic interactions with AS (Ala325, Trp79 and Phe326) and PAS (Tyr329) and CS (His433). Based on ADMET properties, viridiflorene has ability to penetrate BBB and cross GIT while it weak HERG inhibitor (Table 3).

Vomifoliol has anti-AChE activity (Mogana et al. 2014). Vomifoliol has been docked with AChE with BA -7.8 kcal/ mol via hydrogen bonds with Tyr120 in PAS and Gly117 and Gly117 in OH and Ser199 in CS and with several hydrophobic interactions with amino acid residues of PAS (Tyr333 and Asp71) and catalytic triad (Trp83 and His439; SM). Vomifoliol has been docked with BChE (BA of -6.7 kcal/ mol) through hydrogen bonds with three hydroxyl groups of Tyr391 and Thr517 and Asn225 (SM). Based on ADMET results, vomifoliol can be absorbed from GIT and crossed BBB (Table 3).

Cannabispiradienone is a cannabinoid found in cannabis (Pagani et al. 2011). Cannabispiradienone has been docked with AChE (BA -9.2 kcal/mol) via hydrogen bond with Tyr69 residue in PAS and through several hydrophobic interactions with Trp278, Tyr333, Tyr120 and Asp71 in PAS and Phe329 in AS and Phe330 in AP (SM). It has been docked with BChE with BA -8.2 kcal/mol via hydrogen bonding with Asp67 residues of PAS and with hydroxyl group of Thr117 (SM) and through hydrophobic interactions with amino acid residues of AS (Trp79) and CS (His433). The ADMET properties showed it can be crossed BBB and GIT and distributed into mitochondria (Table 3).

Cannabispiran is a cannabinoid found in cannabis (Pagani et al. 2011). In present study, it docked with AChE through hydrogen bond with Asp71 and Tyr69 and through hydrophobic interactions with Tyr333, Trp278 and Tyr120 residues of PAS and Phe329 residues of AS and Phe330 residues of AP (SM). Cannabispiran has a BA – 8.2 kcal/mol

with BChE via hydrogen bonding with Asp67 residues of PAS and with hydroxyl group of Thr117 (SM) and through hydrophobic interactions with amino acid residues of AS (Trp79) and CS (His433). The ADMET properties showed cannabispiran can be crossed BBB and GIT and distributed into mitochondria (Table 3).

Cannabispirol is another cannabinoid found in cannabis (Pagani et al. 2011). In this study, cannabispirol has been docked with AChE with BA (-9.2 kcal/mol) through hydrogen bonding with Tyr120 and several hydrophobic interactions with amino acid residues of PAS (Tyr333), AS (Trp83 and Phe329), AP (Phe330) and CS (His439) and OH (Gly117; Fig. 2). In this line, cannabispirol docked with BChE with BA -8.1 kcal/mol (Table 2) through hydrogen bonds with Trp79 of AS and through hydrophobic interactions with amino acid residues in PAS (Tyr329) and AS (Ala325 and Tyr125) and CS (His433; SM). Based on ADMET properties, it has ability to penetrate BBB and cross GIT while it weak HERG inhibitor (Table 3).

Cannabioxepane is a cannabinoid found in cannabis (Pagani et al. 2011). The present study showed cannabioxepane has been hydrophobically docked with AChE with BA of – 10.4 kcal/mol through Tyr333, Tyr69, Tyr120, Asp71 and Trp278 residues of PAS and Phe330, Phe287 residues of AP and Trp83 and Phe329 of AS and His439 of CS (SM). Cannabioxepane has BA (–9.8 kcal/mol) with BChE via hydrophobic interactions with amino acid residues of PAS (Tyr329 and Asp67) and AS (Ala325 and Trp79) and CS (His433; SM). ADMET properties of cannabioxepane showed that it can go through BBB and absorbed from GIT whilst it is a weak inhibitor of HERG (Table 3).

Isocannabispiradienone is a cannabinoid found in cannabis (Pagani et al. 2011). In this study, isocannabispiradienone docked with AChE with BA –9.4 kcal/mol via hydrogen bonds with AS (Glu198) and with hydroxyl group of Asn84 and via hydrophobic interactions with PAS (Asp71) and CS (His439) and AS (Trp83 and Phe329). It has been docked with BChE (BA –8.3 kcal/mol) via hydrogen bonds with PAS (Asp67 and Tyr329) and AS (Tyr125) and with two hydroxyl groups of Glu194 and Gly112 residues (SM). It can be crossed BBB and GIT and distributed into mitochondria (Table 3).

## Conclusions

Resin of *Cannabis sativa* L. showed an inhibitory effect of 80.00 and 68.00% for AChE and BChE, respectively. The fraction of leaves of *C. sativa* L. showed an inhibitory effect of 52.33 and 49.00% for AChE and BCHE, respectively. In silico results comparable with experimental findings showed that and an array of phytochemicals found in *C. sativa* L. including all cannabinoids, beta-sitosterol, campesterol,

apigenin, alpha-bisabolene,  $\Delta 8$ -THC and  $\Delta 9$ -THC, kaempferol, luteolin, stigmasterol, quercetin, alpha-guaiene, alpha-humulene and caryophyllene oxide have experimentally inhibitory effects for AChE and BChE and would be considered as candidates for designing antiinsectants and/ or anti-AD drugs. In sum, more studies are welcomed to dig deeper mechanisms of anti-ChE properties of *C. sativa* L. and its byproducts like hashish.

Acknowledgement IK and NY conceptualized the idea of this paper and BAH and IK conceived experimental and computational works and wrote the initial draft, and IK and NY critically revised and prepared the final manuscript. This paper originated from MSc thesis of third author submitted to Department of Biology, Faculty of Science, Razi University 67149-67346, Kermanshah, Iran. This study was supported by intramural fund.

#### **Compliance with ethical standards**

**Conflict of interest** No potential conflict of interest was reported by the authors.

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